



>SEQID#1\_SGP037

ATGTCAACAGCTGCCCTTAATTACTTTGGTCAGAAAGTGGTGGGAACCCAGGTGAGAGAGTGTCTGCTAAAGCTCCGCCCTGCTGCAGGACGACA  
GGCGGGTGAACCCACCGTGCCACAGCTCCACTTCAGAGCCTAGGTGTCTCGGTTTGACCCAGATGGTAGTGGGAGTCCAGCTACCTGGGACAA  
TTTGGGATCTGGGATAACCGCATTTGATGAGCCAAATCTGCTGCCACCCAGCAATTAAGTATGGCAAGCCAAATGCCCAAATCAGCTTGGAAATG  
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GCAACAGTAGCCCTATTGCGAGATGGTATTGAACCTGTTGTAGCCAGTGTGGGACAGCCGGCTATTTTGTGTAGAAAGGAAACCCCATGA  
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TAAATTACATCATGCTGATGACAGCTTCCCTGGTCTCCACACAGATGGAAATTAACCTTCATGGTGAATAGTCAAGAGATTGTGACTTTGTCAATC  
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TGGTCCCTGGGAAAATATAAGAACTCTGAAATCAACTTCTCATTCAGCAGAAAGCTTTGCCCTCCAGTGGACGATGGGCCTGA

Fig. 1

>SEQID#2\_SGP037  
MSTAALITLVRSGGNQVRRRVLLSSRLQDDRRVPTICHSTSEPRCSRFPDGGSPATWDFGIWDNRIDEPILLPSIKYKPIPKISLENVGCASQIG  
KRKENEDRFDAQLTDEVLYTAVYDGHGCPAAADFCHTHMEKCTMDLLPKEKNLETLTLAFLHDKAFSSHARLSADATLLTSGTTATVALLRDGIEL  
VVASVGDRAILCRKGKPMKLTIDHTPERKDEKERKKCGGFVAVNSLQPHVNGRLAMTRSIGDLDLKTSGVIAEPETKRKLHHADDSFLVLTIDGI  
NFMVNSQEICDFVNQCHDPNEAAHA VTEQAIQYGTEDNSTAVVFPFGAWGKYKNSEINFSSFSFASGRWA

Fig. 2